

CYR61  
CEF10  
FISP12  
CTGF  
Nov

M--SSSTFRTLAVATLLHL--TRLALST-CPAAC--HCPLE-APKCAPGVGLVRDGGCGCKVCAK  
M--GSAGARP-ALAAALLCL--ARLALGSPCAVC--QCPAA-APQCAPGVGLVPDGGCGCKVCAK  
M--LASVAGPISLA-LVLLALCTRPATGODCSAQC--QCAAEAPHCPAGVSLVDGGCGCCRVCAC  
M--TAASMGVPRVAFVLLALCSRPAVGONCSGPC--RCPDEPAPRPCAGVSLVDGGCGCCRVCAC  
METGGGGLPVLLLLLLLRPCVSGREAAACPRCGGRCPAEP-PRCAPGVPAVLDDGGCGCLVCAR

58  
58  
61  
62  
65

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QLNEDCSKTOPCDHTKGLECNFGASSTALKGICRAQSEGRPCYNSRIYONGESFQPNCKHQCTCI  
QLNEDCSRTOPCDHTKGLECNFGASPAATNGICRAQSEGRPCYNSKIYONGESFQPNCKHQCTCI  
QLGELCTERDPCDPHKGLFCDFGSPANRKIGVCTAK-DGAPCVFGGSVYRSGESFQSSCKYQCTCL  
QLGELCTERDPCDPHKGLFCDFGSPANRKIGVCTAK-DGAPCIFGGTVYRSGESFQSSCKYQCTCL  
ORGESCSPLLPCESGGLYCDRGPEDGGGAGICMVL-EGDNCVFDGMIYRNGETFQPSCKYQCTCR

124  
124  
126  
127  
130

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DGAVGCIPLCPOELSLPNLGCNPRLVKVSGQCCCEWVCEDSIK-DSLDD-QDDLGLDASEVEL  
DGAVGCIPLCPOELSLPNLGCPSRPLVKVPGQCCCEWVCEDESKDALEELEGFFSKEFGLDASEGEL  
DGAVGCVPLCSMDVRLSPDCPFRRVKLPKCKCKEWVCEPKDRTAVGP-----ALAAAYRLED  
DGAVGCMPLCSMDVRLSPDCPFRRVKLPKCKCEEWVCEPKDQTVVGP-----ALAAAYRLED  
DGOIGCLPRCNLGLLLPGPDCPFPRKIEVPGECKWVCDPRDEVLLGGF-----AMAAAYRQEA

188  
190  
185  
186  
189

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FIGURE 1 (1 of 2)

